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Sequence 20241, A Sequence 13471, Ap Sequence 18737, A Sequence 19341, A Sequence 19241, A Sequence 2513, Appl Sequence 2713, Appl Sequence 20174, A Sequence 20174, A Sequence 228, Appl Sequence 228, Appl Sequence 228, Appl Sequence 4976, Ap Sequence 476, Ap Sequence 476, Ap Sequence 8766, Ap Sequence 8766, Ap Sequence 8766, Ap Sequence 2383, Appl Sequence 3253, A
                                                                                                                                                                    Pebruary 16, 2003, 22:05:25 ; Search time 51.0802 Seconds
(Without alignments)
13999.354 Million cell updates/sec
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14: /cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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) US-09-864-761-18737

) US-09-263-959-1

) US-09-864-761-19241

) US-09-864-761-19241

) US-09-864-761-2513

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0 US-09-815-242-8766
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                 nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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Match I
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20 40 2.8 430 10 US-09-815-242-4448 Sequence 2332, Ap 2.8 1455 10 US-09-815-242-4448 Sequence 8394, Ap 2.8 1406 2.8 1476 10 US-09-815-242-8394 Sequence 8394, Ap 2.8 1908 10 US-09-737-178-80 Sequence 50, Appl 2.8 1908 10 US-09-737-178-80 Sequence 50, Appl 2.9 2031 10 US-09-737-178-80 Sequence 84, Appl 2.9 2031 10 US-09-737-178-80 Sequence 84, Appl 2.9 39-6 2.8 996 10 US-09-815-242-875 Sequence 143, Appl 3.9 39-6 2.8 996 10 US-09-815-24857 Sequence 10113, Appl 3.9 39-6 2.8 996 10 US-09-815-24857 Sequence 10113, Appl 3.9 39-6 2.8 640681 10 US-09-815-24857 Sequence 7, Appl 3.9 5 2.8 640681 10 US-09-815-24857 Sequence 7, Appl 3.9 5 2.8 640681 10 US-09-815-24857 Sequence 1, Appl 3.9 5 2.8 640681 10 US-09-915-24857 Sequence 1, Appl 3.9 5 2.8 640681 10 US-09-916-1 Sequence 1, Appl 3.9 5 2.8 640681 10 US-09-916-1 Sequence 1, Appl 3.9 2 2.8 155074 10 US-09-916-1 Sequence 1, Appl 3.9 2 2.8 155074 10 US-09-916-23-40 Sequence 1, Appl 3.9 2 2.8 155074 10 US-09-916-1 Sequence 1, Appl 4.9 38-6 2.7 3666 10 US-09-137-531-1 Sequence 1, Appl 4.9 38-6 2.7 3666 10 US-09-137-531-1 Sequence 1, Appl 4.9 38-6 2.7 3666 10 US-09-137-531-1 Sequence 1, Appl 4.9 38-6 2.7 4197 10 US-09-137-531-8 Sequence 4, Appl 4.9 38-6 2.7 4197 10 US-09-137-531-8 Sequence 4, Appl 4.9 38-6 2.7 4197 10 US-09-137-531-8 Sequence 4, Appl 4.9 38-6 2.7 4197 10 US-09-137-531-8 Sequence 4, Appl 4.9 38-6 2.7 4197 10 US-09-137-531-8 Sequence 4, Appl 4.9 38-6 2.7 513509 9 US-09-137-531-8 Sequence 4, Appl 4.9 38-6 2.7 513509 9 US-09-137-531-8 Sequence 4, Appl 4.9 38-6 2.7 513509 9 US-09-137-531-8 Sequence 4, Appl 4.9 45

ALIGNMENTS

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Sequence 2021, Application US/09864761
September 2021, David R.
APPLICANT: Henzel, David R.
APPLICANT: Henzel, David R.
APPLICANT: Comp. Members US/09/864 761
TITLE OF INVENTION: HUMBER: US/09/864 761
CURRENT APPLICATION NUMBER: US/09/864 761
CURRENT RILNKO DATE: 2001-05-23
PRIOR FILINKO DATE: 2001-05-26
PRIOR FILINKO DATE: 2000-05-26
PRIOR PRIOR PRILOR NUMBER: US/09/632,366
PRIOR FILINKO DATE: 2000-10-30
PRIOR PRILOR PRILOR NUMBER: US/09/632,366
PRIOR FILINKO DATE: 2000-10-30
PRIOR PRILOR PRILOR NUMBER: US/09/00-67
PRIOR FILINKO DATE: 2001-01-30
PRIOR PRILOR PRILOR NUMBER: PCT/US01/0066
PRIOR APPLICATION NUMBER: PCT/US01/0066
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OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 11
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 8.6
OTHER INFORMATION: EXPRESSED IN HELAT, SIGNAL = 9
OTHER INFORMATION: EXPRESSED IN HEL100, SIGNAL = 12
OTHER INFORMATION: EXPRESSED IN HEL100, SIGNAL = 11
OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 10
OTHER INFORMATION: EXPRESSED IN BOULT LIVER, SIGNAL = 8.4
OTHER INFORMATION: EXPRESSED IN BOULT LIVER, SIGNAL = 10
OTHER INFORMATION: EXPRESSED IN BOULT LIVER, SIGNAL = 10
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 10
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 12
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Pred. No. 4.5e-05;
0; Mismatches 650; Indels
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2001-01-29
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annowax Sequence Listing Engine vers. 1.1
SEQ ID NO 20241
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Matches 463; Conservative
                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
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APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES UT
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Aeconica -X-1
CURRENT APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR PELING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/203,366
PRIOR PELING DATE: 2000-00-05-26
PRIOR PELING DATE: 2000-00-04-03
PRIOR PLING DATE: 2000-00-04-07
PRIOR PELING DATE: 2000-00-09-27
PRIOR PELING DATE: 2000-00-09-27
PRIOR PELING DATE: 2001-01-30
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patent No. US20020048763A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 2001-01-30
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4.3%; Score 60; DB 10; Length 1973;
Best Local Similarity 41.5%; Pred. No. 4.9e-05;
Matches 463; Conservative 0; Mismatches 650; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N: MAP TO ACO06547.9

N: EXPRESSED IN LUNG, SIGNAL = 17

N: EXPRESSED IN HELA, SIGNAL = 11

N: EXPRESSED IN HELA, SIGNAL = 8.6

N: EXPRESSED IN HERAT, SIGNAL = 9

N: EXPRESSED IN HELLO, SIGNAL = 12

N: EXPRESSED IN BT474, SIGNAL = 11

N: EXPRESSED IN FETAL LIVER, SIGNAL = 10

N: EXPRESSED IN BONE MARROW, SIGNAL = 8.4

N: EXPRESSED IN BONE MARROW, SIGNAL = 10

N: EXPRESSED IN BONE MARROW, SIGNAL = 10
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2010-10-30
PRIOR PLICATION NUMBER: PCT/US01/00661
PRIOR PLICATION NUMBER: PCT/US01/00661
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR PLICATION NUMBER: PCT/US01/00670
PRIOR PLICATION NUMBER: US 60/234,687
PRIOR PLICATION NUMBER: US 60/24,687
PRIOR PLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR PILING DATE: 2001-01-29
NUMBER: OF SEQ ID NOS: 49117
SEQ ID NO 3471
LENGRIH: 1973
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo saplens
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Sequence 18737, Application US/09864761

Patent No. US20020048763a1

GENERAL INFORMATION:

APPLICANT: Rank, David R.

APPLICANT: Rank, David R.

APPLICANT: Hancal, David R.

CURRENT APPLICATION HUMBER: US 60/180,312

PRIOR PLING DATE: 2000-05-26

PRIOR PLING DATE: 2000-06-03

PRIOR PLICATION NUMBER: US 60/236,356

PRIOR PLICATION NUMBER: US 60/236,356

PRIOR PLICATION NUMBER: CT/US01/0066

PRIOR PLICATION NUMBER: CT/US01/0066

PRIOR PLICATION NUMBER: PCT/US01/0066

PRIOR PLICATION NUMBER: PCT/US01/0066

PRIOR PLICATION NUMBER: PCT/US01/0066

PRIOR FLING DATE: 2001-01-30

PRIOR FLING DATE: 2001-01-30

PRIOR PLICATION NUMBER: PCT/US01/0066

PRIOR FLING DATE: 2001-01-30

PRIOR FLING DATE: 2001-01-30
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TGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTGTCCGGCAATTAAACC 656
                                                                                                                                                                       TGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGGTAGCACAAAACAC 776
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                                                                                                                           TGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATAACCGCATAATGTAACGT
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US-09-864-761-18737/c
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CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: MCMASters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/POCKET NUMBER: 9200;
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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Matches 363; Conservative
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                                                                                                                                                                                                                                                                                           STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              491 TACCGATGCTGAAGCAGGTGTTGATGCAGGTGCTGAAAAAGGTGTTGGTGCAGGTGCAGA 432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            222 IGCTGGTGCTTAACCAAATCCACCTGCTACTGCTAATTTAGTCACATAATGTAACGTTAA 281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                402 TAGTACATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGC 461
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: MAP TO AL035457.11
OTHER INFORMATION: EXPRESSED IN PEACENTA, SIGNAL = 1.2
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 0.85
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.2
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 4.6
OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 4.6
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.93
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.95
OTHER INFORMATION: HIT: U01287.1, EVALUE 1.308+00
US-09-864-761-18737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.7%; Score 52.2; DB 10; Length 46.2%; Pred. No. 0.002; Live 0; Mismatches 203; Indels
                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR PPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers: 1.1
SEQ ID NO 18737
DR APPLICATION NUMBER: PCT/US01/00665
DR FILING DATE: 2001-01-30
DR APPLICATION NUMBER: PCT/US01/00668
DR FILING DATE: 2001-01-30
DR FILING DATE: 2001-01-30
DR FILING DATE: 2001-01-30
DR APPLICATION NUMBER: PCT/US01/00663
DR APPLICATION NUMBER: PCT/US01/00662
                                                                                                                                                                                                             FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00661
arting DATE: 2001-01-30
                                                                                                                                                                                                                                                                                        FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
                                                                                                                                                                                                                                                                                                                                               FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
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Patent No. US20020150891A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              522 AGTAACTACTGATTATG 538
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Matches 174; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-263-959-1
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GENERAL INFORMATION:

APPLICANT: Hood, Leroy E.

APPLICANT: Rowen, Lee

APPLICANT: Koop, Ben F.

TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH NUMBER OF SEQUENCES: 1279
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAATGCCCTGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACA
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Pred. No. 1.1;
0; Mismatches 484;
                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
                                                                                                                                                                                    ADDRESSEE: Seed and Berry LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
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288 IGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTGT 347
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OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 19

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 34

OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 34

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 7.1

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 28

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 25

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 25

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 25

OTHER INFORMATION: EXPRESSED IN HEARY, SIGNAL = 25

OTHER INFORMATION: EXPRESSED IN HEARY, SIGNAL = 18

OTHER INFORMATION: EXPRESSED IN HEARY, SIGNAL = 18

OTHER INFORMATION: EXPRESSED IN HEARY, SIGNAL = 21

OTHER INFORMATION: EXPRESSED IN HEARY, SIGNAL = 21
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                       PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
EENGTH: 1075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 45.8; DB 10;
Pred. No. 0.098;
0; Mismatches 402;
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                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-864-761-19241
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APPLICANT: RAIL ADVIG R.
APPLICANT: RAIL AGAINED: ADVIG R.
TITLE OF INVERTION: HUMAN CENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR FILE OF INVERTION: HUMAN CENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR FILE OF INVERTION NUMBER: US 60/203 ADVIG R.
FILE REFERENCE: 2000-05-23
PRIOR APPLICATION NUMBER: US 60/203 ASVIG REAL APPLICATION NUMBER: PCT/USO1/00667
PRIOR PELICATION NUMBER: PCT/USO1/00669
PRIOR FILING DATE: 2001-01-30
PRIOR FILICATION NUMBER: PCT/USO1/00661
PRIOR PRICK APPLICATION NUMBER: PCT/USO1/00662
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                                                                                                                                                                                                                                                                        13341 TCTCTGACAAGTACTGCTGATGCCACTTAGTACTACTGTACTTATTGCCACTACTTCT 13400
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640 TGTCCGGCAATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATA
                                                                                                         700 ACCGCATAATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAAT
                                                                                                                                                   13281 TCTCCTACAAGTACTACTGTGAGTACTATTGCTACGTTCCCATTTCAGTGACTCCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 19241, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                288 TGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTGT 347
                                                                                                                                                                                                                                                                                                                         348 TAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAAATTTTAATGCAGGTGCTAGTAC 407
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                                                                                                                                                   Length 1403;
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APPLICANT: Lodes, Michael J.
APPLICANT: Houghton, Raymond L.
APPLICANT: HOUGHTON, Raymond L.
APPLICANT: MCNEILL PAITICIAE D.
APPLICANT: MCNEILL BAITICIAE D.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DIAGNOSIS
TITLE OF INVENTION: AND TREATMENT OF EHRLICHIA INFECTION
FILE REFERENCE: 2011.4397.
CURRENT APPLICATION NUMBER: US/09/798,042
CURRENT FILING DATE: 2001-03-02
NUMBER OF SEQ ID NOS: 108
                                                                                                                                                                                                              Indels
25
18
21
                                                                                                                                                   Score 45.8; DB 10;
Pred. No. 0.11;
0; Mismatches 402;
         IN HBL100, SIGNAL = IN HEART, SIGNAL = IN BRAIN, SIGNAL =
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1226 regregregregretreregregregre 1254
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                                                                                                                                                            Ouery Match 3.3%;
Best Local Similarity 41.7%;
Matches 287; Conservative
         CTHER INFORMATION: EXPRESSED CHER INFORMATION: EXPRESSED CTHER INFORMATION: EXPRESSED US-09-864-761-2513
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                         N: MAP TO AL078472.1

N: EXPRESSED IN BONE MARROW, SIGNAL = 27

N: EXPRESSED IN LUNG, SIGNAL = 19

N: EXPRESSED IN BT474, SIGNAL = 34

N: EXPRESSED IN FETAL LIVER, SIGNAL = 7.1

N: EXPRESSED IN ADULT LIVER, SIGNAL = 7.8

N: EXPRESSED IN HELA, SIGNAL = 43

N: EXPRESSED IN PLACENTA, SIGNAL = 43

N: EXPRESSED IN PLACENTA, SIGNAL = 20
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SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 2513
LENGTH: 1403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: GENE EXPRESSION ANALYSIS CURRENT APPLICATION NUMBER: US/09/864,761 CURRENT PILING DATE: 2001-05-23 CURRENT PILING DATE: 2001-05-24 CURRENT FILING DATE: 2001-05-26 CURRENT PRILING DATE: 2000-05-26 PRIOR PRILING DATE: 2000-05-26 PRIOR PRILING DATE: 2000-06-26 PRIOR FILING DATE: 2000-06-36 PRIOR FILING DATE: 2000-06-36 PRIOR FILING DATE: 2000-06-36 PRIOR FILING DATE: 2000-06-36 PRIOR FILING DATE: 2000-00-07 PRIOR FILING DATE: 2000-06-27 PRIOR FILING DATE: 2001-07-37 PRIOR FILING DATE: 2001-07-37 PRIOR FILING DATE: 2001-07-30 PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: PCT/USO1/00664 PRIOR FILING DATE: 2001-01-30 PRIOR FILING PRIOR PRIO
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PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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PRIOR PELING DATE: 2001-01-30
PRIOR PELING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR PELING DATE: 2001-01-30
PRIOR PELING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PELING DATE: 2000-09-21
PRIOR PELING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
PRIOR PELING DATE: 2000-06-30
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR FILING DATE: 2001-01-30
                                                                                                                          TGCAATTGCTAGTGGAGCAACTAATTATG 976
                                                                                                                                                               Sequence 2513, Application US/09864761 patent No. US20020048763A1
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
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ORGANISM: Homo sapiens
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OTHER INFORMATION: M
OTHER INFORMATION: E:
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US-09-864-761-2513
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TELEFAX:
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Patent No. US20020064535A1
GENERAL INFORMATION:
APPLICANT: Reed, Steven G.
APPLICANT: Lodes, Michael J.
APPLICANT: Houghton, Raymond
TITLE OF INVENTION: THERAPY OF EHRLICHIA INFECTION
NUMBER OF SEQUENCES: 73
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                12;
                                                                                                                     Length 2120;
                                                                                                              Score 43.4; DB 10; Length 2
Pred. No. 0.5;
0; Mismatches 336; Indels
              FastSEQ for Windows Version 3.0
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                                                                                                                          Best Local Similarity 44.3
Matches 277; Conservative
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Washington
                                                     TYPE: DNA
ORGANISM: Ehrlichia
SOFTWARE: Fasts
SEQ ID NO 95
                                                                               US-09-798-042-95
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US-09-159-469-39
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979 CAGAGATTACTTTTTTGCGCTACATCAGCATTAGCTTCTACAGATACTTCAGACTTTA 1038
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                                                SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/159,469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
3.1%; Score 43.4; DB 10;
Best Local Similarity 44.3%; Pred. No. 0.5;
Matches 277; Conservative 0; Mismatches 336;
                                                                                                                                                                                                                                                                                                                                        31,392
ER: 210121.439C2
                                                                                                                                                                                       CLASSIFICATION: OPPLIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNMBER: 09/106,582
FILING DATE: 29-JUN-1998
ATTORNEY/ACEDIT INFORMATION:
REGISTRATION NUMBER: 31,392
     COMPUTER: Floppy disk COMPUTER: IBM PC COMPALIBLE OPERATING SYSTEM: PC-DOS/MS-IRPRE: Patenty-
                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGTH: 2129 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                    206-682-6031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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1210 CAGCTACTACAGGTACTTGCGCTAC 1234

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884 AAGCCACTGCAGGTGGTGCCGCTAC 908

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FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 43.4; DB 10; Length 2129; Pred. No. 0.5;
                                                                                                                                                                          APPLICANT: Reed, Steven G.
APPLICANT: Reed, Steven G.
APPLICANT: Lodes, Michael J.
APPLICANT: Lodes, Michael J.
APPLICANT: Houghton, Raymond L.
APPLICANT: MCNEILL, Patricia D.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DIAGNOSIS
TITLE OF INVENTION: AND TREATMENT OF EHRLICHIA INFECTION
FILLE REFERENCE: 21012.1.439C7
CURRENT APPLICATION NUMBER: US/09/798,042
CURRENT FILLING DATE: 2001-03-02
SOFTWARE: FASTSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches 336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    884 AAGCCACTGCAGGTGGTGCCGCTAC 908
1210 CAGCTACTACAGGTACTTGCGCTAC 1234
                                                                                                              Sequence 39, Application US/09798042
Patent No. US20020068343Al
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
CORGANISM: Ehrlichia Sp.
US-09-798-042-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Fast
SEQ ID NO 39
LENGTH: 2129
                                                                                   RESULT 9
US-09-798-042-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     277;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Matches
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PAPLICANT: Chen, Weeshed

TITLE OF INVENTION: HURAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO

TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRY
FILE REPERENCE: Account Cark.

CHERRY TAPLICATION NUMBER: US/09/864,761

CHERRY TAPLICATION NUMBER: US/09/864,761

PRIOR PAPLICATION NUMBER: US/09/864,765

PRIOR PAPLICATION NUMBER: US/09/864,766

PRIOR PALICA DAME: 2000-05-26

PRIOR PALICA DAME: 2000-05-36

PRIOR PALICA DAME: 2000-05-36

PRIOR PALICA DAME: 2000-06-03

PRIOR PALICA DAME: 2001-01-30

PRIOR PALICANION NUMBER: 2001-01-30

PRIOR PAPILCANION NUMBER: 2001-01-30

PRIOR PRIOR PRIOR PAPILCANION NUMBER: 2001-01-30

PRIOR PRIOR PAPILCANION NUMBER: 2001-01-30

PRIOR PRIOR PAPILCANION NUMBER: 2001-01-30

PRIOR PLINK DAME: 2001-01-30

PRIOR PRIOR PAPILCANION NUMBER: 2001-01-30

PRIOR PRIOR PAPILCANION NUMBER: 2001-01-30

PRIOR PAPILCANI
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N: EXPRESSED IN BONE MARROW, SIGNAL = 1.3
N: EXPRESSED IN LUNG, SIGNAL = 1.2
N: EXPRESSED IN HELA, SIGNAL = 1.4
N: EXPRESSED IN HELA, SIGNAL = 1.4
N: EXPRESSED IN HELLOO, SIGNAL = 1.7
N: EXPRESSED IN PLACENTA, SIGNAL = 1.5
N: EXPRESSED IN HEART, SIGNAL = 1.5
N: EXPRESSED IN HEART, SIGNAL = 1.5
N: EXPRESSED IN HEART, SIGNAL = 1.5
N: EXPRESSED IN FETAL LIVER, SIGNAL = 1.5
N: EXPRESSED IN FETAL LIVER, SIGNAL = 1.5
N: EXPRESSED IN FETAL LIVER, SIGNAL = 1.5
Sequence 20174, Application US/09864761 Patent No. US20020048763A1
                                                                                  GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTHER INFORMATION: E:
COTHER INFORMATION: E:
US-09-864-761-20174
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us-09-497-967-3.rnpb

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Sequence 31, Application US/10024623

Fublication No US20020187524A1

GENERAL INFORMATION:
TITLE INFORMATION: 8099, 46455, 54414, 53763, 67076, 67102, 44181,
TITLE OF INVENTION: 8099, 46455, 54414, 53763, 67076, 67102, 44181,
TITLE OF INVENTION: 8099, 46455, 54414, 53763, 67076, 67102, 44181,
TITLE OF INVENTION: 8099, 46455, 54414, 53763, 67076, 67102, 44181,
TITLE OF INVENTION: USE THEREOF
TITLE OF INVENTION: USE THEREOF
TITLE OF INVENTION: USE THEREOF
TITLE OF INVENTION: USE 00.12-17
PRIOR APPLICATION NUMBER: US 60/256,588
PRIOR FILING DATE: 2000-12-18
SPRIOR FILING DATE: 2000-12-21
NUMBER OF SEQ ID NOS: 40
                                                  1014 TAGAATTGGTAATAAACAAACACTACGTACTCGTAGTGGAAAATCTATAGGTGCTAAGGT 1073
                                                                                                                             1074 ACATTATTATTATGATTTAAGTACTATTGATCCTGCAGAAGAAATAGAATTACAAACTAT 1133
                                                                                                                                                                                               990 AGAATGTCTAAATTGTGCTGCTAACTTTTATTTTGATGGTAATAATTTCTAGGCAGGAAG 1049
                                                                                                                                                                                                                                                                            1254 TGTACCCTCTACATCTTTATCAGTTATATTCCTGCAAATACAACAATTCCTTTGGTGG 1313
              TAATAATGCTCCTAATTTCAATCCAGGTAATAGTACATGCCTACCTTGCCCAGCAAATAA 869
                                                                                                                                                                      930 TGCATGCCCTGATGGTACTGCAATTGCTAGTGGAGCAACTAATTATGTAATATTATAAAC 989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4692 TCTTTTTCTAATCCATAATCTGATCATAAATTACACTAATACATTTCTTTTAAAACCT 4633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4632 CAGGATCCTTGTGTTTTGACTATGTATAAGTTTGATGTAAAATAGTAATCCTTTTGTT 4573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GCTAATTGTCCTGTTGGAACTGAAACTAACACAGCCGGATAAGTTGATGATCTAGGAACT 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.0%; Score 42.6; DB 9; 52.5%; Pred. No. 2.3;
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                                                                                                                                                                                                                                                                                                                                                                                                             1110 TACTGCTACTTTAATTGCATAATGTGCCCTTGA 1142
                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 52.5
Matches 116; Conservative
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LENGTH: 25002
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Sequence 1, Application US/09970477

Faten No. US2020127545A1

GENERAL INFORMATION ASSESSMENTS OF HUMAN PAPILLOMA VIRUS-RELATED DISEASE

TITLE OF INVENTION: ASSESSMENTS OF HUMAN PAPILLOMA VIRUS-RELATED DISEASE
CURRENT APPLICATION NUMBER: US/09/970,477

PRIOR PLING DATE: 1908-12-11

PRIOR APPLICATION NUMBER: U.S. 60/082,167

PRIOR PLING DATE: 1998-04-17

PRIOR PLING DATE: 1998-04-17

PRIOR FILING DATE: 1998-04-17

PRIOR FILING DATE: 1998-12-12

PRIOR FILING DATE: 1998-12-12

PRIOR FILING DATE: 1998-12-12

SOFTWARE: PALENTING VINDER: U.S. 60/069,426

NUMBER OF SEQ ID NOS: 5

SOFTWARE: PALENTIN VINDER: 2.1

SEQ ID NO : 2.1
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                                                                                                                                                                                                                                                                                                                                                                                             966 AACTAATTATGTAATATTATAAACAGAATGTCTAAATTGTGCTGCTAACTTTTATTTGA 1025
                                                                        726 TGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGGTAGCACAAAACACTGAATGTAC 785
                                                                                                                                                                                                                                                            262 TGATGGTGATGATGGTGGTGATTACGATGGTGATGCTGCTGCTGCTGCTGGTGGTGATGC 203
                                                                                                    322 CGATGGTGATGGTGATGATGATGGTGATGCTGCTGATGGTGTTGATGGCGATGCTGC 263
                                                                                                                                                                                                                                   846 ATGCCTACCTTGCCCAGCAATAAAGATTATGGTGCTGAAGCCACTGCAGGTGGTGCCGC 905
                                                                                                                                                                                                                                                                                                                906 TACTITAGCCAAATAATGTAATATTGCATGCCCTGATGGTACTGCAATTGCTAGTGGAGC 965
                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                  894 TACATTATATTTTTCTAGTAATGATAATAGTATTAATATATAGCTCCAGATCCTGACTTTTT 953
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                633 CACACCTIGICCGGCAATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGC 692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1026 TGGTAATAATTTCTAGGCAGGAAGTAGTAGATGCAAAGCATGTCCAGCAAATAAAGTT 1083
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     786 TAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTAATTTCAATCCAGGTAATAGTAC
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        Length 439;
Score 42.8; DB 10; Length
Pred. No. 0.35;
0; Mismatches 197; Indels
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Pred. No. 0.65;
0; Mismatches 284;
Query Match
3.0%;
Best Local Similarity 45.0%;
Matches 161; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 3.0%;
Best Local Similarity 44.1%;
Matches 226; Conservative
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US-09-970-477-1
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US-09-970-477-1
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RESULT 13

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Sequence 3346, Application US/09938842A

Sequence 3346, Application US/09938842A

Sequence 3346, Application US/09938842A

Sequence 3346, Application US/09938842A

Setent No. USSO012016

APPLICANT: Harper, Jeff

APPLICANT: Wang, Xun

APPLICANT: Wang, Xun

APPLICANT: Wang, Xun

TITLE OF INVENTION: SAME, AND METHODS OF USE

FILE REFERENCE: SCRIP1300-3

CURRENT APPLICATION NUMBER: US/09/938,842A

CURRENT FILING DATE: 2001-08-24

PRIOR FILING DATE: 2001-08-24

PRIOR APPLICATION NUMBER: US 60/227,866

PRIOR APPLICATION NUMBER: US 60/227,866

PRIOR APPLICATION NUMBER: US 60/227,866

PRIOR PILING DATE: 2001-01-16

PRIOR APPLICATION NUMBER: US 60/200,0111

PRIOR FILING DATE: 2001-01-16

SEQ ID NOS: 5379

SEQ ID NOS: 5379
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  578 ITTACTATAATGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACAC 637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   705 ATAATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGGT 764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  765 AGCACAAAACACTGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTAA 824
                                                                                                                                           525 AACTACTGATTATGTTAGATCATTCACAGAATGTGTTAAATGTAGACTTAACTTTTACTA 584
TGTTAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAG 404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               112 TAGTGGCGGTGGTGCCCATGGTGATGGTGATGGTGATGGTGTTGATAGCTACGGTGATAA 53
                                                645 GGCAATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATAACCGC
                                                                                                                                                                                                                 TACCATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGGAGT
                                                                                                                                                                                                                                                                                                                                                                              352 TAGTGATGGTGATAATGGTGATCATGTGATGGTAGTGGTGGTGCTGCTGATGGTGATGG
                                                                                                                                                                                                                                                                                                                                                                                                                                   585 TAATGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTGTCC
                                                                                                        405 TACATGCACAGGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGCCGC
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2.9%; Score 41.4; DB 9;
Best Local Similarity 54.2%; Pred. No. 1.4;
Matches 84; Conservative 0; Mismatches 71;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          825 TTTCAATCCAGGTAATAGTACATGCC 850
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                   345
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                                                                               PREMERIAL INFORMATION:

PREPLICANT: Penn, SHITTON G.

APPLICANT: Penn, SHITTON G.

APPLICANT: Penn, SHITTON G.

APPLICANT: Hancel, David K.

APPLICANT: Hancel, David K.

APPLICANT: Chen, Wenshed

TITLE OF INVERTION: HURN GROWE-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEPUL FOR TITLE OP INVERTION: GROWE SPRESSION ANALYSIS BY WICKOARRAY CURRENT FILE APPLICANTON NUMBER: 2001-05-23

PRIOR FILID ADDRESSION GROWER GOVERN GROWER GOVERN GOVERN GOVERN GROWER GOVERN GOV
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42.7%; Pred. No. 0.61;
tive 0; Mismatches 290; Indels
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N: EXPRESED IN FETAL LIVER, SIGNAL = 5.2

N: EXPRESED IN HELLO, SIGNAL = 8.2

N: EXPRESED IN BRAIN, SIGNAL = 4.7

N: EXPRESED IN HEART, SIGNAL = 4.3

N: EXPRESED IN ADULT LIVER, SIGNAL = 5.6

N: EXPRESED IN LOUG, SIGNAL = 6.2

N: EXPRESED IN LOUG, SIGNAL = 6.2

N: EXPRESED IN BONE MARROW, SIGNAL = 7.8

N: EXPRESED IN BONE MARROW, SIGNAL = 7.8
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                                                   Sequence 228, Application US/09864761
Patent No. US20020048763A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTHER INFORMATION: M. OTHER INFORMATION: E. US-09-864-761-228
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Best Local Similarity
Matches 216; Conserv
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339 AGAATGTGTTAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGG 398
                                                                                                                                                                                                                                                                                                                                           379 ATATGGTGATAATGGTGATGGTGATGATGATGATGGTAATCATGGTGATGATGG 320
                                                                                                                                                                                                                                                                                                                                                                                                     399 TGCTAGTACATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAA 458
                                                                                                                                                                                                                                                                                                                                                                                                                                                      319 TGGTGATTATGGTAGGGATGGTGATGATAATGGTGATGGTGAGGTGAGGTGATGG 260
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                                                                                                                                                                                                                                               ö
                                                                                                                                                                        2.9%; Score 41; DB 10; Length 489; 50.2%; Pred. No. 1; tive 0; Mismatches 100; Indels
                                                                   - 5.9
                        CTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 5.9 OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL US-09-864-761-4976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Search completed: February 17, 2003, 01:59:09 Job time : 209.08 secs
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Best Local Similarity 50.2%
Matches 101; Conservative
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SOURCE AND APPLICATION US/09864761

SEQUENCE AND APPLICATION US/09864761

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SEQUENCE AND APPLICATION APPLICATION US/09864761

SEQUENCE AND APPLICATION SERVE EXPRESSION ANALYSIS BY MICROARRY

FILLE REPERBECK: Accomica-X: IS/09/0644.761

CURRENT PAPLICATION NUMBER: US/09/0644.761

CURRENT PAPLICATION NUMBER: US/09/0644.761

CURRENT PAPLICATION NUMBER: US/09/0644.761

CURRENT PAPLICATION NUMBER: US/09/064.761

PRIOR FILLING DATE: 2000-09-27

PRIOR FILLING DATE: 2000-09-27

PRIOR FILLING DATE: 2000-09-27

PRIOR PELICATION NUMBER: US/09/0645

PRIOR APPLICATION NUMBER: PCT/08/01/0066

PRIOR APPLICATION NUMBER: PCT/0
OTHER INFORMATION: MAP TO AL031076.1

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 5.1

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 5.5

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 5.8

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 6

OTHER INFORMATION: EXPRESSED IN PLACEMIA, SIGNAL = 5.1

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 5.1

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 5.1
                                                                                  638 CTTGTCCGGCAATTAAACCTGCTAATGTTGCTTAA 672
                                                                                                                                   657 TITGAAAATTAGTTACTTCATTTACAATTTTGTAA 691
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